



SEQUENCE LISTING

<110> Stratagene

<120> Compositions and methods utilizing stable reported cell lines for detection of CHOP-dependent signal

<130> 25436/1490

<140> 09/637,550

<141> 2001-08-11

<150> US 60/150,942

<151> 1999-08-26

<160> 8

<170> PatentIn version 3.0

<210> 1

<211> 6

<212> DNA

<213> Artificial

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<223> contiguous residues

<220>

<221> misc_feature

<222> (1)..(6)

<223> contiguous residues

<400> 1

ggatcc

6

<210> 2

<211> 6

<212> DNA

<213> Artificial

<220>

<223> conserved sequence

<220>

<221> misc_feature

<222> (1)..(6)

<223> conserved sequence

n at position 3 is any purine residue.

n at position 4 is any pyrimidine residue

<400> 2

ggnncc

6

B4

<210> 3
<211> 99
<212> PRT
<213> Unknown

<220>
<223> mammalian c-Jun sequence

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<222> (1)..(99)
<223> transactivation domain

<400> 3

Met Ala Ala Glu Ser Leu Pro Phe Thr Leu Glu Thr Val Ser Ser Trp
1 5 10 15

Glu Leu Glu Ala Trp Tyr Glu Asp Leu Gln Glu Val Leu Ser Ser Asp
20 25 30

Glu Ile Gly Gly Thr Tyr Ile Ser Ser Pro Gly Asn Glu Glu Glu
35 40 45

Ser Lys Thr Phe Thr Thr Leu Asp Pro Ala Ser Leu Ala Trp Leu Thr
50 55 60

Glu Glu Pro Gly Pro Thr Glu Val Thr Arg Thr Ser Gln Ser Pro Arg
65 70 75 80

Ser Pro Asp Ser Ser Gln Ser Ser Met Ala Gln Glu Glu Glu Glu Glu
85 90 95

Glu Gln Gly

<210> 4
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<213> Unknown

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cggagtactg tcctccg

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<212> DNA
<213> Unknown

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gtcggagtac tgtcctccga gcggagtact gtcctccgag cggagtactg tcctccgagc 60
ggagtactgt cctccgagcg gagtactgtc ctccgagcgg agactctaga ggggtatata 120
atggatcccc gggtagcgag ctccaattc 149

B⁴
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<211> 31
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<222> (1)..(31)
<223> plasmid sequence

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cagcttgga ttccgtact gttggtaat g 31

<210> 7
<211> 74
<212> DNA
<213> Artificial

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<223> plasmid sequence

<220>
<221> misc_feature
<222> (1)..(74)
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gtatcgccgg gatccgccg ggctggaatt ctagaagctt ctgcagagct cggtagcaga 60
tcttgaataa gtag 74

<210> 8
<211> 21
<212> PRT
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<220>
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B4
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<222> (1)..(21)
<223> selected fusion activation domain

<400> 8

Val Ser Pro Gly Ser Gly Arg Ala Gly Ile Leu Glu Ala Ser Ala Glu
1 5 10 15

Leu Gly Thr Arg Ser
20
